

2113 45



OIPE

**RAW SEQUENCE LISTING**  
**PATENT APPLICATION: US/09/902,903**

**DATE: 02/19/2002**  
**TIME: 14:08:10**

**Input Set : D:\CRF sequence listing.txt**  
**Output Set: N:\CRF3\02192002\I902903.raw**

3 <110> APPLICANT: Genentech, Inc.  
 4 Ashkenazi, Avi  
 5 Botstein, David  
 6 Desnoyers, Luc  
 7 Eaton, Dan L.  
 8 Ferrara, Napoleone  
 9 Filvaroff, Ellen  
 10 Fong, Sherman  
 11 Gao, Wei-Qiang  
 12 Gerber, Hanspeter  
 13 Gerritsen, Mary E.  
 14 Goddard, A.  
 15 Godowski, Paul J.  
 16 Grimaldi, Christopher J.  
 17 Gurney, Austin L.  
 18 Hillan, Kenneth, J.  
 19 Kljavin, Ivar J.  
 20 Mather, Jennie P.  
 21 Pan, James  
 22 Paoni, Nicholas F.  
 23 Roy, Margaret Ann  
 24 Stewart, Timothy A.  
 25 Tumas, Daniel  
 26 Williams, P. Mickey  
 27 Wood, William, I.  
 29 <120> TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic  
 30 Acids Encoding the Same  
 32 <130> FILE REFERENCE: 10466-14  
 C--> 34 <140> CURRENT APPLICATION NUMBER: US/09/902,903  
 C--> 35 <141> CURRENT FILING DATE: 2001-07-10  
 37 <150> PRIOR APPLICATION NUMBER: PCT/US00/04414  
 38 <151> PRIOR FILING DATE: 2000-02-22  
 40 <150> PRIOR APPLICATION NUMBER: US 60/143,048  
 41 <151> PRIOR FILING DATE: 1999-07-07  
 43 <150> PRIOR APPLICATION NUMBER: US 60/145,698  
 44 <151> PRIOR FILING DATE: 1999-07-26  
 46 <150> PRIOR APPLICATION NUMBER: US 60/146,222  
 47 <151> PRIOR FILING DATE: 1999-07-28  
 49 <150> PRIOR APPLICATION NUMBER: PCT/US99/20594  
 50 <151> PRIOR FILING DATE: 1999-09-08  
 52 <150> PRIOR APPLICATION NUMBER: PCT/US99/20944  
 53 <151> PRIOR FILING DATE: 1999-09-13  
 55 <150> PRIOR APPLICATION NUMBER: PCT/US99/21090

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PATENT APPLICATION: US/09/902,903

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TIME: 14:08:11

Input Set : D:\CRF sequence listing.txt  
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56 <151> PRIOR FILING DATE: 1999-09-15  
 58 <150> PRIOR APPLICATION NUMBER: PCT/US99/21547  
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 62 <151> PRIOR FILING DATE: 1999-10-05  
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 65 <151> PRIOR FILING DATE: 1999-11-29  
 67 <150> PRIOR APPLICATION NUMBER: PCT/US99/28313  
 68 <151> PRIOR FILING DATE: 1999-11-30  
 70 <150> PRIOR APPLICATION NUMBER: PCT/US99/28564  
 71 <151> PRIOR FILING DATE: 1999-12-02  
 73 <150> PRIOR APPLICATION NUMBER: PCT/US99/28565  
 74 <151> PRIOR FILING DATE: 1999-12-02  
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 77 <151> PRIOR FILING DATE: 1999-12-16  
 79 <150> PRIOR APPLICATION NUMBER: PCT/US99/30911  
 80 <151> PRIOR FILING DATE: 1999-12-20  
 82 <150> PRIOR APPLICATION NUMBER: PCT/US99/30999  
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 84 <150> PRIOR APPLICATION NUMBER: PCT/US00/00219  
 85 <151> PRIOR FILING DATE: 2000-01-05  
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 91 <211> LENGTH: 1825  
 92 <212> TYPE: DNA  
 93 <213> ORGANISM: Homo sapiens  
 95 <400> SEQUENCE: 1  
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 97 gaccacgcg tccgggccgg agcagcacgg ccgcaggacc tggagctccg gtcgtgtctt 120  
 98 cccgcagcgc taccgcgcatt ggcctgcgg cgccggccg cgctggggct cctggcgctt 180  
 99 ctgtctgtgc tgccgcgcgc gccggaggcc gccaagaagc cgacgcgcctg ccaccgggtgc 240  
 100 cgggggctgg tggacaagtt taaccagggg atggtgacca ccgcaaagaa gaactttggc 300  
 101 ggccggaaaca cggctggga gaaaaagacg ctgttcaagt acgagtccag cgagattcgc 360  
 102 ctgctggaga tcctggaggg gctgtgcgag agcagcgcact tcgaatgc当地 tcagatgcta 420  
 103 gaggcgcagg aggacacact ggaggcctgg tggctgcgc tgaagagcga atatcctgac 480  
 104 ttattcgagt ggtttgtgt gaagacactg aaagtgtgt gctctccagg aacctacgtt 540  
 105 cccgactgtc tcgcattgcca gggcgatcc cagaggccct gcaagcgggaa tggccactgc 600  
 106 agcggagatg ggagcagaca gggcgacggg tcctggccgt gcccacatggg gtaccaggc 660  
 107 ccgcgtgtca ctgactgcatt ggacggctac ttcaagctgc当地 tccggaaacga gaccacagc 720  
 108 atctgcacag cctgtgacga tcctgtcaag acgtgtctgg gcctgaccaa cagagactgc 780  
 109 ggcgagtgtg aagtgggtgt ggtgtggac gagggcgccct gtgtggatgt ggacgagtgt 840  
 110 gcggccgagc cgcctccctg cagcgctgc当地 cagttctgtta agaacgc当地 cggctccctac 900  
 111 acgtgc当地 agtgtgactc cagctgtgtg ggctgc当地 gagggaaaggccc aggaaaactgt 960  
 112 aaagagtgta tctctggctt cgc当地 gagggag cacgacagtg gtgc当地 gagatgt ggacgagtgc 1020  
 113 tcacttagcag aaaaaacctg tgtgaggaaa aacgaaaact gctacaatac tccaggc当地 1080  
 114 tacgtctgtg tgtgtctgtc当地 cggcttc当地 gaaacggaaag atgc当地 ctgtgt gccc当地 cggc当地 1140  
 115 gaggctgaag ccacagaagg agaaaagcccg acacagctgc当地 cctcccgca agacctgtaa 1200  
 116 tgtgccc当地 ac gatgc当地 ggac gggaggct gcctgctc当地 taacgggtta 1260  
 117 gatgc当地 ggccgtc tcctgc当地 agt gacagc当地 ggccg gggaggct gcctgctc当地 taacgggtta 1320

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118 ttctcatttg tcccttaaac agctgcattt cttgggttgtt cttaaacaga cttgtatatt 1380
119 ttgatacagt tctttgtaat aaaattgacc atttaggta atcaggagga aaaaaaaaaa 1440
120 aaaaaaaaaaa aaaggcgcc cgcgactcta gagtcgacct gcagaagctt ggcgcctatg 1500
121 gcccaacttg ttatgcag cttataatgg ttacaataaa agcaatagca tcacaaattt 1560
122 cacaataaaa gcatttttt cactgcattc tagtgtggt ttgtccaaac tcataatgt 1620
123 atcttatcat gtctggatcg ggaattaatt cggcgacgca ccatggcctg aaataacctc 1680
124 tgaaagagga acttggtagt gtaccttcg aggccgaaag aaccagctgt ggaatgtgtg 1740
125 tcaatgggg tgtggaaagt ccccaggctc cccagcaggc agaagtatgc aagcatgcat 1800
126 ctcaattagt cagcaaccca gttt 1825
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129 <211> LENGTH: 353
130 <212> TYPE: PRT
131 <213> ORGANISM: Homo sapiens
133 <400> SEQUENCE: 2
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135 1 5 10 15
137 Leu Leu Pro Pro Ala Pro Glu Ala Ala Lys Lys Pro Thr Pro Cys His
138 20 25 30
140 Arg Cys Arg Gly Leu Val Asp Lys Phe Asn Gln Gly Met Val Asp Thr
141 35 40 45
143 Ala Lys Lys Asn Phe Gly Gly Asn Thr Ala Trp Glu Glu Lys Thr
144 50 55 60
146 Leu Ser Lys Tyr Glu Ser Ser Glu Ile Arg Leu Leu Glu Ile Leu Glu
147 65 70 75 80
149 Gly Leu Cys Glu Ser Ser Asp Phe Glu Cys Asn Gln Met Leu Glu Ala
150 85 90 95
152 Gln Glu Glu His Leu Glu Ala Trp Trp Leu Gln Leu Lys Ser Glu Tyr
153 100 105 110
155 Pro Asp Leu Phe Glu Trp Phe Cys Val Lys Thr Leu Lys Val Cys Cys
156 115 120 125
158 Ser Pro Gly Thr Tyr Gly Pro Asp Cys Leu Ala Cys Gln Gly Gly Ser
159 130 135 140
161 Gln Arg Pro Cys Ser Gly Asn Gly His Cys Ser Gly Asp Gly Ser Arg
162 145 150 155 160
164 Gln Gly Asp Gly Ser Cys Arg Cys His Met Gly Tyr Gln Gly Pro Leu
165 165 170 175
167 Cys Thr Asp Cys Met Asp Gly Tyr Phe Ser Ser Leu Arg Asn Glu Thr
168 180 185 190
170 His Ser Ile Cys Thr Ala Cys Asp Glu Ser Cys Lys Thr Cys Ser Gly
171 195 200 205
173 Leu Thr Asn Arg Asp Cys Gly Glu Cys Glu Val Gly Trp Val Leu Asp
174 210 215 220
176 Glu Gly Ala Cys Val Asp Val Asp Glu Cys Ala Ala Glu Pro Pro Pro
177 225 230 235 240
179 Cys Ser Ala Ala Gln Phe Cys Lys Asn Ala Asn Gly Ser Tyr Thr Cys
180 245 250 255
182 Glu Glu Cys Asp Ser Ser Cys Val Gly Cys Thr Gly Glu Gly Pro Gly
183 260 265 270
185 Asn Cys Lys Glu Cys Ile Ser Gly Tyr Ala Arg Glu His Gln Cys

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|     |   |     |     |     |
|-----|---|-----|-----|-----|
| 186 | 275   | 280 | 285 |     |
| 188 | Ala Asp Val Asp Glu Cys Ser Leu Ala Glu Lys Thr Cys Val Arg Lys                 |     |     |     |
| 189 | 290   | 295 | 300 |     |
| 191 | Asn Glu Asn Cys Tyr Asn Thr Pro Gly Ser Tyr Val Cys Val Cys Pro                 |     |     |     |
| 192 | 305   | 310 | 315 | 320 |
| 194 | Asp Gly Phe Glu Glu Thr Glu Asp Ala Cys Val Pro Pro Ala Glu Ala                 |     |     |     |
| 195 | 325   | 330 | 335 |     |
| 197 | Glu Ala Thr Glu Gly Glu Ser Pro Thr Gln Leu Pro Ser Arg Glu Asp                 |     |     |     |
| 198 | 340   | 345 | 350 |     |
| 200 | Leu   |     |     |     |
| 203 | <210> SEQ ID NO: 3  |     |     |     |
| 204 | <211> LENGTH: 2206  |     |     |     |
| 205 | <212> TYPE: DNA   |     |     |     |
| 206 | <213> ORGANISM: Homo sapiens  |     |     |     |
| 208 | <400> SEQUENCE: 3   |     |     |     |
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| 210 | tcgacacctga cccacgcgtc cgccaggccg ggaggcgacg cgcacccagccg tctaaaacggg 120       |     |     |     |
| 211 | aacagccctg gctgaggggag ctgcagcgca gcagagtatac tgacggcgcc aggttgcgtta 180        |     |     |     |
| 212 | gggtcgccac gaggagttt cccggcgacg aggaggctt gaggcagcatg gcccgaggaga 240           |     |     |     |
| 213 | gcgccttccc tgccggcccg ctctggctt ggagcatcct cctgtgcctg ctggcactgc 300            |     |     |     |
| 214 | gggcggaggc cgggcggccg caggaggaga gcctgtaccc atggatcgat gctcaccagg 360           |     |     |     |
| 215 | caagagtaact cataggattt gaagaagata tcctgattgt ttcagagggg aaaatggcac 420          |     |     |     |
| 216 | cttttacaca tgatttcaga aaagcgcaac agagaatgcc agctattcct gtcaatatcc 480           |     |     |     |
| 217 | attccatgaa ttttacctgg caagctgcag ggcaggcaga atacttctat gaattcctgt 540           |     |     |     |
| 218 | ccttgcgtc cctggataaa ggcatcatgg cagatccaaac cgtcaatgtc cctctgctgg 600           |     |     |     |
| 219 | gaacagtgcc tcacaaggca tcagttgttc aagtgggtt cccatgttggg gaaaaacagg 660           |     |     |     |
| 220 | atgggggtggc agcatttgaa gtggatgtga ttgttatgaa ttctgaaggc aacaccattc 720          |     |     |     |
| 221 | tccaaacacc tcaaaatgtc atcttcttta aaacatgtca acaagctgag tgcccaggcg 780           |     |     |     |
| 222 | ggtgccgaaa tggaggcttt tgtaatgaaa gacgcatctg cgagtgtcct gatgggttcc 840           |     |     |     |
| 223 | acggacactca ctgtgagaaa gccccttgc ccccacgtatg tatgaatggg ggactttgtg 900          |     |     |     |
| 224 | tgactcctgg tttctgcattc tgcccacctg gattctatgg agtgaactgt gacaaagcaa 960          |     |     |     |
| 225 | actgctcaac cacctgtttt aatggaggga cctgtttcta ccctggaaaa tgtatttgcc 1020          |     |     |     |
| 226 | ctccaggact agaggaggag cagtgtgaaa tcagcaaattt cccacaaccc tgcgaaatg 1080          |     |     |     |
| 227 | gaggttaatg cattgtaaa agcaaatgtt aagtgttccaa aggttaccatgg ggagacctct 1140        |     |     |     |
| 228 | gttcaagcc ttgtctgcgtg cttggctgtg gtgcacatgg aacctgcccattt gaacccaaca 1200       |     |     |     |
| 229 | aatgccaatg tcaagaaggt tggcatggaa gacactgcaaa taaaaggtac gaagccagcc 1260         |     |     |     |
| 230 | tcatacatgc cctggggcca gcaggcgccc agctcaggca gcacacgcct tcacttaaaa 1320          |     |     |     |
| 231 | aggccgagga gcggcgggat ccacctgtat ccaattacat ctggtaact ccgacatctg 1380           |     |     |     |
| 232 | aaacgtttta agttcacca agttcatagc ctttgcattt ctttcatgtt gatggatgttc 1440          |     |     |     |
| 233 | aaataatgtt cattacactt aagaataactg gcctgtatattt tattagctt attataaattc 1500       |     |     |     |
| 234 | actgagctga tattttacttcc ttgttcaatgt acgtctgttag catgtatggta 1560                |     |     |     |
| 235 | tagattttct tttttcaatgtt ctttgggaca gattttatattt tatgtcaattt gatcagggtt 1620     |     |     |     |
| 236 | aaattttcaatgtt tttttcaatgtt ctttgggaca gattttatattt tatgtcaattt gatcagggtt 1680 |     |     |     |
| 237 | ggggcggggg acatcagaaa gttttttttt gggaaaaatggc gttttttttt gttttttttt 1740        |     |     |     |
| 238 | atgggtcagt taatgttggaa gttttttttt gttttttttt gttttttttt gttttttttt 1800         |     |     |     |
| 239 | ttttttttt tttttttttt gttttttttt tttttttttt gttttttttt gttttttttt 1860           |     |     |     |
| 240 | ttttttttt tttttttttt gttttttttt gttttttttt gttttttttt gttttttttt 1920           |     |     |     |
| 241 | ttttttttt tttttttttt gttttttttt gttttttttt gttttttttt gttttttttt 1980           |     |     |     |
| 242 | ttttttttt tttttttttt gttttttttt gttttttttt gttttttttt gttttttttt 2040           |     |     |     |

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243 ttttatactg tttgtatgta taaaataaaag gtgctgctt agtttttgg aaaaaaaaaa 2100
244 aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa gggccggccgc gactctagag tcgacctgca 2160
245 gaagcttggc cgccatggcc caacttgtt attgcagctt ataatg 2206
247 <210> SEQ ID NO: 4
248 <211> LENGTH: 379
249 <212> TYPE: PRT
250 <213> ORGANISM: Homo sapiens
252 <400> SEQUENCE: 4
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257 20 25 30
259 Glu Glu Ser Leu Tyr Leu Trp Ile Asp Ala His Gln Ala Arg Val Leu
260 35 40 45
262 Ile Gly Phe Glu Glu Asp Ile Leu Ile Val Ser Glu Gly Lys Met Ala
263 50 55 60
265 Pro Phe Thr His Asp Phe Arg Lys Ala Gln Gln Arg Met Pro Ala Ile
266 65 70 75 80
268 Pro Val Asn Ile His Ser Met Asn Phe Thr Trp Gln Ala Ala Gly Gln
269 85 90 95
271 Ala Glu Tyr Phe Tyr Glu Phe Leu Ser Leu Arg Ser Leu Asp Lys Gly
272 100 105 110
274 Ile Met Ala Asp Pro Thr Val Asn Val Pro Leu Leu Gly Thr Val Pro
275 115 120 125
277 His Lys Ala Ser Val Val Gln Val Gly Phe Pro Cys Leu Gly Lys Gln
278 130 135 140
280 Asp Gly Val Ala Ala Phe Glu Val Asp Val Ile Val Met Asn Ser Glu
281 145 150 155 160
283 Gly Asn Thr Ile Leu Gln Thr Pro Gln Asn Ala Ile Phe Phe Lys Thr
284 165 170 175
286 Cys Gln Gln Ala Glu Cys Pro Gly Gly Cys Arg Asn Gly Gly Phe Cys
287 180 185 190
289 Asn Glu Arg Arg Ile Cys Glu Cys Pro Asp Gly Phe His Gly Pro His
290 195 200 205
292 Cys Glu Lys Ala Leu Cys Thr Pro Arg Cys Met Asn Gly Gly Leu Cys
293 210 215 220
295 Val Thr Pro Gly Phe Cys Ile Cys Pro Pro Gly Phe Tyr Gly Val Asn
296 225 230 235 240
298 Cys Asp Lys Ala Asn Cys Ser Thr Thr Cys Phe Asn Gly Gly Thr Cys
299 245 250 255
301 Phe Tyr Pro Gly Lys Cys Ile Cys Pro Pro Gly Leu Glu Gly Glu Gln
302 260 265 270
304 Cys Glu Ile Ser Lys Cys Pro Gln Pro Cys Arg Asn Gly Gly Lys Cys
305 275 280 285
307 Ile Gly Lys Ser Lys Cys Lys Cys Ser Lys Gly Tyr Gln Gly Asp Leu
308 290 295 300
310 Cys Ser Lys Pro Val Cys Glu Pro Gly Cys Gly Ala His Gly Thr Cys
311 305 310 315 320
313 His Glu Pro Asn Lys Cys Gln Cys Glu Gly Trp His Gly Arg His

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→ Use of n and/or Xaa has been detected in the Sequence Listing.  
Review the Sequence Listing to insure a corresponding  
explanation is presented in the <220> to <223> fields of  
each sequence using n or Xaa.

VERIFICATION SUMMARY  
PATENT APPLICATION: US/09/902,903

DATE: 02/19/2002  
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Input Set : D:\CRF sequence listing.txt  
Output Set: N:\CRF3\02192002\I902903.raw

L:34 M:270 C: Current Application Number differs, Replaced Current Application Number  
L:35 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:516 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13  
L:517 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13  
L:518 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13  
L:519 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13  
L:774 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26  
L:1706 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50  
L:3591 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:113  
L:4045 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:131  
L:5349 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:174  
L:5484 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:175  
L:6545 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:206